

SEQUENCE LISTING

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<120> Adeno-associated virus vectors
<130> 875.007US2
<140> US10/054,665
<141> 2002-01-22
<150> US 60/086,166
<151> 1998-05-20
<150> US 09/276,625
<151> 1999-03-25
<160> 14
<170> FastSEQ for Windows Version 4.0
<210> 1
<211> 20
<212> DNA
<213> Adeno-associated virus
<400> 1
cgggggtcgt tgggcggtca
                                                                         20
<210> 2
<211> 19
<212> DNA
<213> Adeno-associated virus
<400> 2
gggcggagcc tatggaaaa
                                                                         19
<210> 3
<211> 505
<212> DNA
<213> Artificial Sequence
<220>
<223> A synthetic consensus sequence
<400> 3
cgggggtcgt tgggcggtca gccaggcggg ccatttaccg taagttatgt aacgactgca
                                                                         60
ggcatgcaag ctcgaattca tcggtagata agtagcatgg cgggttaatc attaactaca
                                                                        120
aggaacccct agtgatggag ttggccactc cctctctgcg cgctcgctcg ctcgctgagg
                                                                        180
ccgggcgacc aaaggtcgcc cgacgcccgg gctttgcccg ggcggcctca gtgagcgagc
                                                                        240
gagegegeag etgegegete getegeteae tgaggeegee egggeaaage eegggegteg
                                                                        300
ggcgaccttt ggtcgcccgg cctcagcgag cgagcgagcg cgcagagagg gagtggccaa
                                                                        360
ctccatcact aggggttcct tgtagttaat gattaacccg ccatgctact tatctacagc
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ttgcatgcat gtgagcaaaa ggccagcaaa aggccaggaa ccgtaaaaag gccgcgttgc

tggcgttttt ccataggctc cgccc

420

480

505

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<210> 4
<211> 272
<212> DNA
<213> AAV circular intermediate, clone p81
<400> 4
gcatgcaagc tgtagataag tagcatggcg ggttaatcat taactacaag gaacccctag
                                                                         60
tgatggagtt ggccactccc tctctgcgcg ctcgctcgct cactgaggcc gggcggccaa
                                                                        120
                                                                        180
aggtcgcccg acgcccgggc tttgcccggg cggcctcagt gagcgagcga gcgcgcagag
                                                                        240
agggagtggc caactccatc actaggggtt ccttgtagtt aatgattaac ccgccatgct
acttatctac cgatgaattc gagcttgcat gc
                                                                        272
<210> 5
<211> 300
<212> DNA
<213> AAV circular intermediate, clone p79
<400> 5
gcatgcaagc tgtagataag tagcatggcg ggttaatcat taactacaag gaacccctag
                                                                         60
tgatggagtt ggccactccc tctctgcgcg ctcgctcgct cactgaggcc gggcgcgcgc
                                                                        120
tegetegete actgaggeeg ggegaecaaa ggtegeeega geeegggett tgeeegggeg
                                                                        180
gcctcagtga gcgagcgcgc gcgcagagag ggagtggcca actccatcac taggggttcc
                                                                        240
ttgtagttaa tgattaaccc gccatgctac ttatctaccg atgaattcga gcttgcatgc
                                                                        300
<210> 6
<211> 272
<212> DNA
<213> AAV circular intermediate, clone p1202
<400> 6
gcatgcaagc tgtagataag tagcatggcg ggttaatcat taactacaag gaacccctag
                                                                         60
tgatggagtt ggccactccc tctctgcgcg ctcgctcgct cactgaggcc gggcgaccaa
                                                                        120
aggtcgcccg acgcccgggc tttggtcgcc cggcctcagt gagcgagcga gcgcgcagag
                                                                       180
agggagtggc caactccatc actaggggtt ccttgtagtt aatgattaac ccgccatgct
                                                                        240
acttatctac cgatgaattc gagcttgcat gc
                                                                       272
<210> 7
<211> 165
<212> DNA
<213> Unknown
<220>
<223> SEQ ID NO:1 of U.S. Patent No. 5,478,745
<400> 7
aggaacccct agtgatggag ttggccactc cctctctgcg cgctcgctcg ctcactgagg
                                                                         60
ccgggcgacc aaaggtcgcc cgacgcccgg gctttgcccg ggcggcctca gtgagcgagc
                                                                       120
gagcgcgcag agagggagtg gccaactcca tcactagggg ttcct
                                                                        165
<210> 8
<211> 282
<212> DNA
<213> rAAV circular intermediate, clone p79
<400> 8
ggcgggccat ttaccgtaag ttatgtggcg actgcaggca tgcaagctcg aattcatcgg
                                                                         60
tagataagta gcatggcggg ttaatcattg cctacaaaga gcccctagtg atggagtggg
                                                                        120
ccactccctc tcttcgccga gcgcgcagag agggagtggc caactccctc actaggggtt
                                                                       180
cctggcagtt aatgattaac ccgccatgct acttatctac agcttgcatg catgtgagca
                                                                        240
aaaggccagc aaaaggccag gaaccgtaaa aaggccgcgt tg
                                                                       282
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<210> 9
<211> 345
<212> DNA
<213> rAAV circular intermediate, clone p80
<400> 9
ggccatttac cgtaagttat gtaacgactg caggcatgca agctcgaatt catcggtaga
                                                                         60
taagtagcat ggcgggttaa tcattaacta caaggaaccc ctagtgatgg agttggccac
                                                                        120
tecetetetg egegeteget egetegetea ggeegggega ecaaaggteg eeegaegeee
                                                                        180
gcccggcctc agcgagcgag cgagcgcgca gagagggagt ggccaactcc atcactaggg
                                                                        240
gttccttgta gttaatgatt aacccgccat gctacttatc tacagcttgc atgcatgtga
                                                                        300
gcaaaaggcc agcaaaaggc caggaaccgt aaaaaggccg cgttg
                                                                        345
<210> 10
<211> 276
<212> DNA
<213> rAAV circular intermediate, clone p81
<400> 10
ggccatttac cgtaagttat gtggcgactg caggcatgca agctcgaatt catcggtaga
                                                                         60
taagtagcat ggcgggttaa tcattgccta caaagagccc ctagtgatgg agcccggcct
                                                                        120
caccgagcga gcgagcgcgc agagagggag tggccaactc catcactagg ggttccttgt
                                                                        180
agttaatgat taacccgcca tgctacttat ctacagcttg catgcatgtg agcaaaaggc
                                                                        240
cagcaaaagg ccaggaaccg taaaaaggcc gcgttg
                                                                        276
<210> 11
<211> 316
<212> DNA
<213> rAAV circular intermediate, clone p86
<400> 11
ggccatttac cgtaagttat gtaacgactg caggcatgca agctcgaatt catcggtaga
                                                                         60
taagtagcat ggcgggttaa tcattaacta caaggaaccc ctagtgatgg agttggccac
                                                                        120
tecetetetg egegeteget egetegetga ggeegeeeeg geeteagega gegagegage
                                                                        180
gcgcagagag ggactggcca actccatcac taggggttcc ttgtagttaa tgattaaccc
                                                                        240
gccatgctac ttatctacag cttgcatgca tgtgagcaaa aggccagcaa aaggccagga
                                                                        300
accgtaaaaa ggccgc
                                                                        316
<210> 12
<211> 208
<212> DNA
<213> rAAV circular intermediate, clone p87
<400> 12
ggccatttac cgtaagttat gtaacgactg caggcatgca agctcgaatt catcggtaga
                                                                         60
taagtagcat ggcgggttac tcattgccta caaagagccc ctagtgatgg aattggaatg
                                                                        120
attcaccctc catgctactt atctacagct tgcatgcatg tgagcaaaag gccagcaaaa
                                                                       180
ggccaggaac cgtaaaaagg ccgcgttg
                                                                       208
<210> 13
<211> 310
<212> DNA
<213> rAAV circular intermediate, clone p88
<400> 13
gccatttacc gtaagttatg taacgactgc aggcatgcaa gctcgaattc atcggtagat
                                                                        60
aagtagcatg gcgggttaat cattgcctac aaagagcccc tagtgatgga gttggccact
                                                                       120
ccctctctgc gcgctcgctc gctgggcccg gcctcagcga gcgagcgagc gcgcagagag
                                                                       180
ggaguggeea actecateae taggggttee ttgtagttaa tgattaacce gecatgetae
                                                                       240
ttatctacag cttgcatgca tgtgagcaaa aggccagcaa aaggccagga accgtaaaaa
                                                                       300
ggccgcgttg
                                                                       310
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<210> 14
<211> 334
<212> DNA
<213> Artificial Sequence
<220>
<223> A synthetic portion of the consensus sequence
<400> 14
gtagataagt agcatggcgg gttaatcatt aactacaagg aacccctagt gatggagttg
                                                                       60
gccactccct ctctgcgcgc tcgctcgctc gctgaggccg ggcgaccaaa ggtcgcccga
                                                                      120
cgcccgggct ttgcccgggc ggcctcagtg agcgagcgag cgcgcagctg cgcgctcgct
                                                                      180
cgctcactga ggccgcccgg gcaaagcccg ggcgtcgggc gacctttggt cgcccggcct
                                                                      240
cagcgagcga gcgagcgcgc agagagggag tggccaactc catcactagg ggttccttgt
                                                                      300
agttaatgat taacccgcca tgctacttat ctac
                                                                      334
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